Reviewer Report

Title: Galaxy mothur Toolset (GmT): a user-friendly application for 16S rRNA gene sequencing analysis using mothur.

Version: Original Submission Date: 4/16/2018

Reviewer name: Xianfeng Chen

Reviewer Comments to Author:

The revised version has addressed some of my concerns including a little more introductions about Galaxy, package installations, large file transfer, etc. Yet, it could not completely convince me without a demonstration how robust this tool is, a plot of running time vs. the number of total reads (starting from uploading fastq files) may help as described in 1.b in my previous comments.

Or, the authors can emphasize the novelty by, for example, involving more downstream analysis tools to make this tool a upgrade version of mothur as described in the last paragraph of my previous comments.

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